Query= SEQ ID NO:1
(618 letters)

	Score	E
Sequences producing significant alignments:	(bits)	Value
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>AC114490.2.1.152428
Length = 152428

Score = 1222 bits (616), Expect = 0.0
Identities = 617/618 (99%)
Strand = Plus / Minus

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Sbjct: 107249  atgggggctgcctttgtcgctagcctccgcagtaacctttcttctgccacttcaaggtca 107190


Query: 61      gagatgaacagcagtgttggggacctgggtgttggcggctgcagcctctgggatgaccct 120
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Query: 361     acgtgtccacctatgcggcggtggtcttcgccgcgctcatcagcgtgtgccgctgcggct 420
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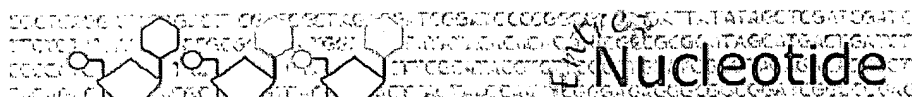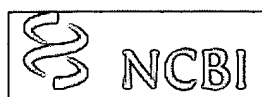
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Query: 481     cgcgcgctgcctgcgccttcgcctggctggcgggcctggcccctccctgcctggagcacc 540
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Query: 541     gctgggcaagctcggggctggcctccgccacggtggccttcgcggccgccttcctgctgg 600
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Query: 601     tgctcgcggccaacgtga 618
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Sbjct: 106649 tgctcgcggccaacgtga 106632
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NCBI

Search [Nucleotide ▼] for [   ] [Go] [Clear]

Limits   Preview/Index   History   Clipboard   Details

[Display] [default ▼] Show: [20 ▼] [Send to] [File ▼]   [Get Subsequence]

☐ **1: AC114490. Homo sapiens chro...[gi:20340495]**   Links

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LOCUS       AC114490              152428 bp    DNA     linear   PRI 30-APR-2002
DEFINITION  Homo sapiens chromosome 1 clone RP11-244H3, complete sequence.
ACCESSION   AC114490 AL354876
VERSION     AC114490.2  GI:20340495
KEYWORDS    HTG.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 152428)
  AUTHORS   Kaul,R.K., Olson,M.V., Zhou,Y., James,R.A., Rouse,G., Wu,Z.,
            Saenphimmachak,C., Phelps,K.A., Buckley,D., Raymond,C. and
            Haugen,E.D.
  TITLE     Direct Submission
  JOURNAL   Unpublished
REFERENCE   2  (bases 1 to 152428)
  AUTHORS   Kaul,R.K., Olson,M.V., Raymond,C. and Haugen,E.D.
  TITLE     Direct Submission
  JOURNAL   Submitted (09-MAR-2002) Genome Center, University of Washington,
            Box 352145, Seattle, WA 98195, USA
REFERENCE   3  (bases 1 to 152428)
  AUTHORS   Kaul,R.K., Olson,M.V., Zhou,Y., James,R.A., Rouse,G., Wu,Z.,
            Saenphimmachak,C., Phelps,K.A., Buckley,D., Raymond,C. and
            Haugen,E.D.
  TITLE     Direct Submission
  JOURNAL   Submitted (30-APR-2002) Genome Center, University of Washington,
            Box 352145, Seattle, WA 98195, USA
COMMENT     On Apr 30, 2002 this sequence version replaced gi:19310309.
            -------------- Genome Center
              Center: University of Washington Genome Center
              Center Code: UWGC
              Web site: http://www.genome.washington.edu
              Contact: uwgchtgs@u.washington.edu
              Drafting Center: SC
            -------------- Project Information
              Center project name: chr-1
              Center clone name: RP11-244H3 (sc0659)
            ------------- Summary Statistics
              Sequencing vector: plasmid; 31% of reads
              Sequencing vector: plasmid; L08752; 69% of reads
              Chemistry: Dye-terminator ET; 89% of reads
              Chemistry: Dye-terminator Big Dye; 11% of reads
              Assembly program: Phrap; version 0.990319
              Consensus quality: 152402 bases at least Q40
              Consensus quality: 152428 bases at least Q30
              Consensus quality: 152428 bases at least Q20
              Insert size: 152428; sum-of-contigs
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